

1 Mark R. Vermeulen [CSBN 115381]  
Law Office of Mark R. Vermeulen  
2 755 Florida Street #4  
San Francisco, CA 94110.2044  
3 Phone: 415.824.7533  
Fax: 415.824.4833  
4 vermeulen@mindspring.com

5 Bicka Barlow [CSBN 178723]  
Law Office of Bicka Barlow  
6 2358 Market St.  
San Francisco, CA 94114  
7 Phone: 415.553.4110  
Fax: 415-553-4110  
8 bickabarlow@sbcglobal.net

9 Attorneys for Defendant  
CHARLES HEARD

11 UNITED STATES DISTRICT COURT  
12 NORTHERN DISTRICT OF CALIFORNIA  
13 SAN FRANCISCO DIVISION

14 UNITED STATES OF AMERICA,

15 Plaintiff,

16 v.

17 CHARLES HEARD,

18 Defendant.

NO. 13-cr-00764-WHO

**DEFENDANT CHARLES HEARD'S  
SUPPLEMENTAL MEMORANDUM RE:  
DNA DAUBERT HEARING**

Hearing Date: May 16, 2017  
Time: 9:00 a.m.  
Judge: William H. Orrick

21 **INTRODUCTION**

22 Defendant Charles Heard, through counsel, files this memorandum in connection with the  
23 DNA *Daubert* hearing scheduled for May 16-17, 2017. The Court previously ordered the hearing  
24 on the admissibility of DNA evidence based on the motion filed by Defendant Adrian Gordon, in  
25 which Mr. Heard joined. *See Defendant Adrian Gordon's Motion/Memorandum To Exclude DNA*  
26 *Test Results And Request For Daubert Hearing* (Dkt. No. 639) and the related *Exhibits* (Dkt. No.  
27 639-1); *Defendant Charles Heard's Joinder In Codefendant Adrian Gordon's Motion To Exclude*  
28

1 *DNA Test Results And Request For A Daubert Hearing* (Dkt. No. 638); *see also* Defendant Esau  
 2 Ferdinand's *Motion to Exclude Evidence of DNA Testing Performed by Serological Research*  
 3 *Institute and Request for Daubert Hearing and Memorandum of Points and Authorities* (Dkt. Nos.  
 4 647 & 648); *Defendant Charles Heard's Joinder in Codefendants' Motions* (including Mr.  
 5 Gordon's and Mr. Ferdinand's DNA/*Daubert* motions) (Dkt. No. 683). The DNA *Daubert*  
 6 hearing will include defendants Heard, Gordon and Ferdinand. The DNA testing at issue for Mr.  
 7 Heard was conducted by Serological Research Institute (SERI), as was the DNA testing at issue  
 8 for Mr. Ferdinand.

9 As stated in the earlier motions, Mr. Heard asserts that: (1) there is no reliable scientific  
 10 basis for this proposed testimony, and thus the testimony is inadmissible under *Daubert v. Merrell*  
 11 *Dow Pharmaceuticals, Inc.*, 509 U.S. 579 (1993) and *Kumho Tire Co. v. Carmichael*, 526 U.S.  
 12 137 (1999); and (2) the testimony is inadmissible under the amendments to Fed.R.Evid. Rule 702  
 13 effective as of Dec. 1, 2000, in that (a) the testimony is not based upon sufficient facts or data, (b)  
 14 the testimony is not the product of reliable principles and methods, and (c) the government's DNA  
 15 technicians have not applied the principles and methods reliably to the facts of the case.<sup>1</sup> Mr.  
 16 Heard also asserts that even if the Court were to find the DNA evidence admissible under *Daubert*  
 17 and Rule 702, the Court should exclude the evidence under Fed.R.Evid. 403 because it is  
 18 substantially more prejudicial, confusing, and misleading than probative.

19 The underlying legal principles and general factual basis regarding Low Copy Number  
 20 (LCN) or Low Template (LT) DNA testing was set forth in Mr. Gordon's moving papers.<sup>2</sup> In this  
 21

22  
 23 <sup>1</sup> While the DNA *Daubert* issues were initiated in this case via pleadings framed as motions to exclude, the  
 24 Government, as the proponent of the DNA evidence, bears the burden of demonstrating that the DNA  
 testing is reliable and that the results are admissible. *Daubert*, 509 U.S. at 592 n.10; *United States v. Orr*,  
 692 F.3d 1079, 1091 (10th Cir. 2012); Fed.R.Evid. 702 advisory committee's note to 2000 amendment.

25 <sup>2</sup> The terms LCN and LT have similar if slightly different meanings to lab analysts. LCN was a term first  
 26 coined in 2000 to apply to testing in which enhanced methods were used to obtain more sensitive results.  
 27 As the test kits themselves became more sensitive and labs began to apply stochastic thresholds, the term  
 LT came into use. LT DNA generally does not require the use of enhancements in the testing; rather, it  
 simply refers to samples with very low levels of DNA. In this motion, the term LCN will be used to refer  
 28 to both LCN and LT testing.

Supplemental Memorandum, Mr. Heard addresses some of the specific issues regarding the testing done in his case, similar to those elucidated in Mr. Gordon's briefing.

### **SUMMARY OF FACTS PERTINENT TO SERI'S TESTING AND ANALYSIS<sup>3</sup>**

The DNA test results from SERI are considered LCN DNA testing for two reasons: 1) The data from the DNA testing of the two samples the government intends to offer fall within the stochastic range of the test; and 2) the lab used two different enhanced methods in conducting the testing, as set forth below. In addition, SERI used a statistical method that is specifically disapproved by the forensic science community.

The evidence at issue are DNA test results from three swabs taken from the left rear door of the vehicle in which two persons were found murdered on August 14, 2008. These swabs, Item 5-1 (left rear door lock) and Item 6-1 (left rear door handle) were obtained and tested by SERI using the Identifiler Plus test kit.<sup>4</sup> Prior to amplifying the sample with Identifiler Plus, the lab quantified the samples. The lab reported that Item 5-1 contained 0.008 ng/ul of total human DNA in a volume of 30 ul, and Item 6-1 contained 0.0028 ng/ul of total human DNA in 30 ul.<sup>5</sup> SERI amplified 0.08 ng of Item 5-1 and 0.25 ng of Item 6-1.<sup>6</sup> The lab used its "ID+29" amplification set up which calls for 29 cycles of amplification.<sup>7</sup> The lab used a 10 second injection and doubled the amount of DNA injected for Item 6-1, and did a 10 second injection of Item 5-1.<sup>8</sup> Both increased injection and doubling of the loaded volume are designed to increase the sensitivity of the test and are listed in the SWGDAM Guidelines for STR Enhanced Detection Methods.<sup>9</sup>

<sup>3</sup> This summary is presented to provide the Court with a basic framework of the basis for this *Daubert* challenge; it is not intended to be comprehensive.

<sup>4</sup> Bates BG078541 – BG078549. The Bates pages cited in this memorandum are filed under seal in an accompanying filing.

<sup>5</sup> Bates BG084110. The abbreviation "ng" stands for nanogram, which is one billionth of a gram ( $10^{-9}$  g.). The abbreviation "ul" stands for microliter, which is one millionth of a liter ( $10^{-6}$  g.).

<sup>6</sup> *Id.*

<sup>7</sup> Bates BG084390; SERI Methods Manual, "STR Typing – Genetic Analyzer" at p. 3 of 17 (effective date 01/11/2011).

<sup>8</sup> Bates BG084279 and BG084283.

<sup>9</sup> SWGDAM is the abbreviation for the Scientific Working Group on DNA Analysis Methods.

1 The lab obtained results indicating that the two samples were complex mixed samples.  
 2 Item 5-1 was a mixture of at least 3 individuals and Item 6-1 was a mixture of at least 4  
 3 individuals.<sup>10</sup> Based on the lab's quantitation records, and an assumption of equal contributors to  
 4 Items 5-1, approximately 0.026 ng or 26 pg<sup>11</sup> of DNA was amplified for each contributor, and for  
 5 Item 6-1, with the same assumption, approximately 0.063 ng or 63 pg of DNA was amplified for  
 6 each contributor.<sup>12</sup> A human cell contains approximately 6 pg of DNA. Thus, for individuals in  
 7 Item 5-1, three equal portions would consist of approximately 4 or 5 human cells worth of DNA  
 8 from each contributor. For Item 6-1, four equal portions would consist of approximately 10  
 9 human cells worth of DNA from each contributor.

10 The lab concluded that both "Charles Heard could be a contributor to item 5-1," and  
 11 "[a]pproximately one person in 12,600 could also be similarly included with respect to Heard.  
 12 Isiah Turner could be a minor contributor to this DNA.<sup>13</sup> Approximately 1 person in 155 could  
 13 also be similarly included with respect to Turner."<sup>14</sup> The lab concluded that as to Item 6-1,  
 14 "Charles Heard could be a contributor to item 6-1," and "[a]pproximately one person in 20 could  
 15 also be similarly included with respect to Heard."<sup>15</sup> The lab also concluded that Isiah Turner and  
 16 Reginald Elmore could be minor contributors to Item 6-1, and that "[a]pproximately one person  
 17 in 18 could also be similarly included with respect to Isiah Turner and approximately one person  
 18 in 2 could also be a minor contributor with respect to Elmore."<sup>16</sup>

19 The match statistic employed by SERI in drawing these numerical conclusions is the  
 20 Combined Probability of Inclusion (CPI). This statistic has been used on mixed samples in which

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21 <sup>10</sup> Bates BG078547.

22 <sup>11</sup> The abbreviation "pg" stands for picogram, which is one trillionth of a gram ( $10^{-12}$  g.)

23 <sup>12</sup> This calculation was done simply to illustrate the very small amounts of DNA present per individual in  
 24 this mixture. Given the complexity of the mixture it is not possible to tell exact ratios, and it is not possible  
 to assess the actual number of individuals in the sample.

25 <sup>13</sup> Bates BG078547. Mr. Turner is one of the persons murdered on Aug. 14, 2008.

26 <sup>14</sup> *Id.*

27 <sup>15</sup> *Id.*

28 <sup>16</sup> *Id.* The Government has indicated that it does not seek to admit into evidence the testing results  
 regarding Mr. Elmore.

1 a single profile cannot be discerned from the mixture, and it estimates the probability that an  
2 individual chosen at random would not be excluded as a potential donor to the evidence sample.

3 The application of a match statistic in DNA testing is a two-step process. First, the analyst  
4 makes a subjective decision regarding whether an individual is included or excluded as a potential  
5 donor. Second, the analyst chooses the loci and alleles or types to include in the statistic and  
6 calculates a statistic. The scientific literature clearly indicates that the use of the CPI method may  
7 not be appropriate under certain circumstance, including where data has dropped out. P. Gill, et  
8 al., *DNA Commission of the International Society of Forensic Genetics: Recommendations on the*  
9 *Interpretation of Mixtures*, 160 Forensic Sci. Int. 90, 95, 101 (2006).

10 In particular it is problematical to apply the [CPI] method when there are  
11 loci which, under the hypothesis being considered of the suspect at hand,  
12 appear to have alleles in category C (alleles not seen due to dropout). We  
13 have seen many instances in which laboratories do just this, usually by  
14 omitting from the RMNE calculation the inconvenient loci. Such a  
15 calculation implies, certainly incorrectly, that among the “random men”  
16 considered for comparison by the calculation only the same loci would be  
17 used for inculcation/exculpation as those being considered for the present  
18 suspect. It fails to acknowledge that choosing the omitted loci is suspect-  
19 centric and therefore prejudicial against the suspect.

20 *Id.* at 91.

21 Additionally, the CPI calculation does not allow for stutter and dropout to be assessed  
22 probabilistically. The DNA Commission recommended that the use of the CPI approach be  
23 restricted to “unambiguous” profiles (mixture profiles in which major and minor contributors can  
24 be clearly distinguished), and described the likelihood ratio statistic as the correct approach. The  
25 Commission also indicated that if the DNA crime stain profile contains low levels of DNA, and if  
26 drop-out is possible, the CPI method is not the appropriate method to use. *Id.* at 91-92.

27 In this instance, the use of the CPI is inappropriate. The samples in both Item 5-1 and Item  
28 6-1 are complex mixtures of at least three or four individuals, respectively; the data is low level (as  
determined by the input quantity of DNA and the amount of data below the stochastic threshold);  
and the lab must invoke allelic drop-out in order to include Mr. Heard in both samples. Because

1 these samples are low level and the lab invokes dropout, Items 5-1 and 6-1 are considered to be  
2 LCN samples.

3       Additionally, SERI's CPI calculations are "suspect-centric": they involve a specific,  
4 known suspect. In fact, for all of the samples tested by SERI it reports out different CPI  
5 calculations for each individual whose reference (or known) sample was tested. This is a clear  
6 violation of Guideline 4.2 of the SWGDAM Interpretation Guidelines for Autosomal STR Typing,  
7 January 2010, which states, in relevant part: "For calculating the CPE [another name for CPI] or  
8 RMP, *any DNA typing results used for statistical analysis must be derived from evidentiary*  
9 *items and not known samples.*" (Emphasis added.) Furthermore, scrutiny of SERI's CPI  
10 calculation shows that the lab does not include alleles below the stochastic threshold in the  
11 calculation, nor does it consider peaks in the stutter position. While on its face the exclusion of  
12 unreliable data from this statistic might be seen as conservative, it actually is not. The impact of  
13 not considering these alleles statistically is that it makes the number of individuals who might be  
14 included in the mixture appear to be fewer. The end result is that the statistic appears to make an  
15 inclusion more rare when, in fact, if that data were considered in the statistic, the resulting CPI  
16 would indicate a more common inclusion. As an example, if the CPI that did not use the low level  
17 alleles was 1 in 100,000, when those alleles were included the CPI could drop to a much lower  
18 number such as 1 in 50,000. The failure to consider this data statistically actually prejudices the  
19 defendant and is misleading in its statement of rarity.

20       The results for both tests fall within the lab's defined stochastic range by peak height. In  
21 this case, in order to include Mr. Heard as a contributor to either Item 5-1 or Item 6-1, the lab must  
22 invoke the phenomenon of allelic dropout at at least one locus for Item 5-1 and at least two loci for  
23 Item 6-1. In addition, as demonstrated by the statistical analysis of both samples, the lab has  
24 invoked a defendant-centered statistic which ignores exculpatory information and is specifically  
25 disapproved by SWGDAM.

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# **SERI'S DNA TESTING AND ANALYSIS IS INADMISSIBLE**

In its role as the gatekeeper under Fed.R.Evid. 702, the Court must “ensure that evidence presented by expert witnesses is *relevant, reliable, and helpful to the jury's evaluation* of such evidence.” *Daubert*, 509 U.S. at 589, 597 (emphasis added). While *Daubert* set forth a non-exclusive list of factors for the Court to consider, those factors are “meant to be helpful, not definitive, and the trial court has discretion to decide how to test an expert’s reliability as well as whether the testimony is reliable, based on the particular circumstances of the particular case.” *City of Pomona v. SQM North America Corp.*, 750 F.3d 1036 (9th Cir. 2014) (quoting *Primiano v. Cook*, 598 F.3d 558, 564 (9th Cir. 2010) (citations and quotation marks omitted)).

This gatekeeping function “requires the judge to assess the reasoning and methodology underlying the expert’s opinion, and determine whether it is both scientifically valid and applicable to a particular set of facts.” *Dodge v. Cotter Corp.*, 328 F.3d 1212, 1221 (10th Cir. 2003). The Supreme Court has made clear that “where [expert] testimony’s factual basis, data, principles, methods, or their application are called sufficiently into question . . . the trial judge must determine whether the testimony has ‘a reliable basis in the knowledge and experience of [the relevant] discipline.’” *Kumho Tire Co.*, 526 U.S. at 149 (quoting *Daubert*, 509 U.S. at 592).

Generally, the district court should focus on an expert’s methodology rather than the conclusions he/she generates. *Daubert*, 509 U.S. at 595. However, an expert’s conclusions are not immune from scrutiny: “A court may conclude that there is simply too great an analytical gap between the data and the opinion proffered.” *General Elec. Co. v. Joiner*, 522 U.S. 136, 146 (1997). “[N]othing in either *Daubert* or the Federal Rules of Evidence requires a district court to admit opinion evidence which is connected to existing data only by the *ipse dixit* of the expert.” *Dodge*, 328 F.3d at 1222.

While a number of cases have addressed the specific question of LCN DNA testing, Mr. Heard urges this Court to adopt the rationale in *United States v. McCluskey*, 954 F.Supp.2d 1224 (D.N.M. 2013). In *McCluskey*, the Court was presented with extensive briefing on many issues related to DNA testing and extensive testimony on the issues involved in LCN DNA testing. As



1 to one sample tested, the Court concluded that under *Daubert*, the tests results were unreliable.  
 2 The *McCluskey* Court's analysis differed from other cases<sup>17</sup> in that the focus for the Court "[was]  
 3 not to establish a definition of 'LCN testing,' but to determine whether the Government's DNA  
 4 results in this case are reliable and admissible," and noted that the laboratory's own protocol  
 5 "declared [the sample in question] to contain too small a quantity to yield the ***normally reliable***  
 6 DNA profile obtained through PCR/STR testing; Item 1B23B, at 215 pg, [as that sample] is below  
 7 the Lab's 'stochastic threshold' and therefore 'in the potential danger zone of unreliable results.'" *Id.*  
 8 at 1278 (emphasis added). The Court expressly recognized that other labs may have differing  
 9 protocols and obtain reliable results with lower quantities of DNA than the lab in *McCluskey*, and  
 10 emphasized the issue in the case:

11 But what is important is that the NMDPS Lab has empirically determined  
 12 that 250 pg is its own stochastic threshold; this means that the NMDPS Lab  
 13 recognizes that, applying its own protocols and using its own  
 14 instrumentation, ***it expects to see stochastic effects that may render the***  
 15 ***results unreliable*** when a sample under 250 pg is tested. The question  
 16 before the Court is whether the Government has carried its burden of  
 17 demonstrating, by a preponderance of the evidence, that the LCN testing by  
 18 the NMDPS Lab in this case is nevertheless reliable.

16 *Id.* at 1279 (emphasis added).

17 The *McCluskey* Court rejected the Government's argument that the fact that other labs  
 18 performed similar testing made the evidence in the case admissible, and focused its analysis on  
 19 whether the testing in *this* case was reliable.<sup>18</sup> *Id.* In determining the reliability of the test  
 20 results, the Court consider the presence of peak height imbalance, the fact that increased injection  
 21 time made the sample ineligible for upload to CODIS, the fact that no replicate testing was done,  
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23 <sup>17</sup> The different approaches and cases are discussed in Mr. Gordon's moving papers. See *Defendant Adrian*  
 24 *Gordon's Memorandum* (Dkt. No. 639) at 18:21-22:16.

25 <sup>18</sup> The *McCluskey* Court also rejected the argument that the holding in *People v. Megnath*, 27 Misc.3d 405,  
 26 898 N.Y.S.2d 408 (N.Y.Sup.Ct. 2010) was persuasive because the lab at issue used different procedures  
 27 and interpretive methods, that there was extensive validation and received certification of its methods, that  
 28 the *Megnath* held extensive hearings and determined the credibility of that testimony, that the *Megnath*  
 court "made the questionable findings that the *Frye* test only applies to novel scientific evidence and that  
 LCN testing was 'not a novel scientific technique' (despite change in procedure and interpretive methods)"  
 and that the Court didn't account for increased stochastic effects." *McCluskey*, 954 F.Supp.2d at 1279-80.



1 and finally, that the analyst “did not present a good explanation, or scientific basis, for her  
2 opinion that the profile on 1B23B was reliable despite the clear stochastic effects of dropout at  
3 four or five loci.” *Id.* at 1283-84. The *McCluskey* Court concluded that the Government had not  
4 met the standard for admissibility of the testing under *Daubert* and Rule 702 and had “not carried  
5 its burden of demonstrating, by a preponderance of the evidence, that LCN testing done by the  
6 NMDPS Lab is reliable and admissible under *Daubert* and Rule 702.” *Id.* at 1287-88.

### 7 CONCLUSION

8 In this case, the DNA evidence proffered by the Government should be excluded under  
9 *Daubert* and Fed.R.Evid. 702 because the methodology used by SERI produces unreliable data  
10 and the conclusion of inclusion by SERI in based on the unreliable data obtained by the lab,  
11 thereby making that conclusion unreliable. Furthermore, SERI applied a statistical analysis of  
12 the data that is specifically disapproved by the forensic community. Finally, even if the Court  
13 were to find the DNA evidence admissible under *Daubert* and Rule 702, the Court should  
14 exclude the evidence under Fed.R.Evid. 403 because it is substantially more prejudicial,  
15 confusing, and misleading than probative.

16 Dated: May 10, 2017

Respectfully submitted,

17  
18 /S/

19 Mark R. Vermeulen  
20 Bicka Barlow  
21 Attorneys for Defendant  
22 CHARLES HEARD  
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